

Figure 1

Polynucleotide sequence of Human BMP2A gene (gi: 179501 and 4557368) (SEQ ID NO:1)
(Coding sequence: 324 – 1514)

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1 ggggacttct tgaacttgca gggagaataa cttgctgcacc ccactttgcg ccggtgcctt
61 tgccccagcg gagcctgctt cgccatctcc gagccccacc gccctccac tcctcggcct
121 tgccccagac tgagacgctg ttcccagcgt gaaaagagag actgcgcggc cggcaccg
181 gagaaggagg aggcaaagaa aaggaacgga cattcggtcc ttgcccagg tcctttgacc
241 agagtttttc catgtggacg ctctttcaat ggacgtgtcc ccgctgtctt cttagacgga
301 ctgcggtctc ctaaaggctg accatggtgg ccgggaccgg ctgtcttcta gcgttgctgc
361 ttccccaggt cctcctgggc ggcgcggtg gcctcgttcc ggagctgggc cgcaggaagt
421 tcgcggcggc gtcgtcggc cgccctcat ccagccctc tgacgaggtc ctgagcgagt
481 tcgagttgcy gctgctcagc atgttcggcc tgaacagag acccaccgcc agcagggacg
541 ccgtggtgcc cccctacatg ctagacctgt atcgcaggca ctcaggtcag ccgggctcac
601 ccgccccaga ccaccggttg gagagggcag ccagccgagc caacactgtg cgcagcttcc
661 accatgaaga atctttggaa gaactaccag aaacgagtgg gaaaacaacc cggagattct
721 tctttaattt aagttctatc cccacggagg agtttatcac ctcagcagag cttcaggttt
781 tccgagaaca gatgcaagat gctttaggaa acaatagcag tttccatcac cgaattaata
841 tttatgaaat cataaaacct gcaacagcca actcgaaatt ccccgtagacc agacttttgg
901 acaccagggtt ggtgaatcag aatgcaagca ggtgggaaag ttttgatgtc acccccgtg
961 tgatgcggtg gactgcacag ggacacgcca accatggatt cgtggtggaa gtggccact
1021 tggaggagaa acaagggtgc tccaagagac atgttaggat aagcaggtct ttgcaccaag
1081 atgaacacag ctggtcacag ataaggccat tgctagtaac ttttggccat gatggaaaag
1141 ggcattcctc ccacaaaaga gaaaaacgtc aagccaaaca caaacagcgg aaacgcctta
1201 agtccagctg taagagacac cttttgtacg tggacttcag tgacgtgggg tggaatgact
1261 ggattgtggc tccccgggg tatcacgct tttactgcca cggagaatgc ccttttctc
1321 tggctgatca tctgaactcc actaatcatg ccattgttca gacgttggtc aactctgtta
1381 actctaagat tcctaaggca tgctgtgtcc cgacagaact cagtgtatc togatgctgt
1441 accttgacga gaatgaaaag gttgtattaa agaactatca ggacatggtt gtggagggtt
1501 gtgggtgtcg ctagtacagc aaaattaaat acataaatat atatata

Figure 2

Polypeptide sequence of human BMP2A (SEQ ID NO:2 = translation of SEQ ID NO:1)

| | | | | | |
|----|-----|------------|------------|-------------|------------|
| 5 | 1 | MVAGTRCLLA | LLLPQVLLGG | AAGLVPELGR | RKFAAASSGR |
| | 41 | PSSQPSDEVL | SEFELRLLSM | FGLKQRPTPS | RDAVVPPYML |
| | 81 | DLYRRHSGQP | GSPAPDHRLE | RAASRANTVR | SFHHEESLEE |
| | 121 | LPETSGKTTR | RFFFNLSIP | TEEFITSAEL | QVFREQMQDA |
| | 161 | LGNNSSFHHR | INIYEIIKPA | TANSKFPVTR | LLDTRLVNQN |
| 10 | 201 | ASRWESFDVT | PAVMRWTAQG | HANHGFEVVEV | AHLEEKQGVV |
| | 241 | KRHVRISRSL | HQDEHSWSQI | RPLLVTFGHD | GKGHPLHKRE |
| | 281 | KRQAKHKQRK | RLKSSCKRHP | LYVDFSDVGW | NDWIVAPPGY |
| | 321 | HAFYCHGECF | FPLADHLNST | NHAIVQTLVN | SVNSKIPKAC |
| | 361 | CVPTELSAIS | MLYLDENEKV | VLKNYQDMVV | EGCGCR |

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Figure 3
Results of a loss of function validation experiment

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